

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/519,505

CRF Edit Date: 2/24/06
Edited by: JKR

___ **Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line**

___ **Corrected the SEQ ID NO. Sequence numbers edited were:**

___ **Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:**

☒ Deleted: ☒ Invalid beginning/end-of-file text ; ___ page numbers

___ **Inserted mandatory headings/numeric identifiers, specifically:**

___ **Moved responses to same line as heading/numeric identifier, specifically:**

___ **Other:**

**Raw Sequence Listing before editing,
for reference only**

BEST AVAILABLE COPY



PC1

RAW SEQUENCE LISTING

DATE: 02/24/2006

PATENT APPLICATION: US/10/519,505

TIME: 12:33:16

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\02242006\J519505.raw

3 <110> APPLICANT: Aarhus Universitet
 5 <120> TITLE OF INVENTION: Disease risk estimating method using sequence polymorphisms in a
 specific
 6 region of chromosome 19
 8 <130> FILE REFERENCE: P 687 PC00
 -> 10 <140> CURRENT APPLICATION NUMBER: US/10/519,505
 -> 10 <141> CURRENT FILING DATE: 2004-12-27
 10 <160> NUMBER OF SEQ ID NOS: 172
 12 <170> SOFTWARE: PatentIn version 3.1
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 16 <212> TYPE: DNA
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Does Not Comply
 Corrected Diskette Needed
 (P5.6) ↗

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<212> DNA
.. <213> Artificial sequence

<220>
<223> Probe

<400> 172
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21

??
??
??
??
1

deleted

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/519,505

DATE: 02/24/2006
TIME: 12:33:17

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\02242006\J519505.raw

valid Line Length:

rules require that a line not exceed 72 characters in length. This includes spaces.

g#:1; Line(s) 5

VERIFICATION SUMMARY

DATE: 02/24/2006

PATENT APPLICATION: US/10/519,505

TIME: 12:33:17

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\02242006\J519505.raw

10 M:270 C: Current Application Number differs, Replaced Current Application No
10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
1603 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 ✓
1605 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:172
1605 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 ✓
1607 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:172
1607 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 ✓
1609 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:172
1609 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 ✓
1611 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:172